AMENDMENT

In the Claims:

Please replace the presently pending claims with the following claims:

(Twice amended) An isolated 125P5C8 protein comprising the sequence of SEQ 1. ID NO: 2. (Amended) A 125P5C8 protein, wherein the 125P5C8 protein has at least 6 2. contiguous amino acids of an amino acid sequence shown in SEQ ID NO: 2. 3. (Amended) The 125P5C8 protein of claim 2, wherein 125P5C8 protein has at least 15 contiguous amino acids of an amino acid sequence shown in SEQ ID NO: 2. (Amended) The 125P5C8 protein of claim 3, wherein the 125P5C8 protein is at 4. least 30 contiguous anino acids of an amino acid sequence shown in SEQ ID NO: 2. 7. (Amended) An 125P5C8 protein of claim 1 that further comprises at least one conservative substitution. 8. (Twice amended) An 125P5C8 protein of claim 1 that comprises an epitope that induces a specific antibody response. (Twice amended) An isolated 125P5C8 protein of claim 1 that has an amino acid 14. sequence which is exactly that of an amino acid sequence encoded by a polynucleotide selected

- from the group consisting of:
 - (a) a polynucleotide consisting of the sequence as shown in SEQ ID NO: 1,
- a polynucled tide consisting of the sequence as shown in SEQ ID NO: 1, from (b) nucleotide residue number 82\through nucleotide residue number 696,
- a polynucleotide that encodes a 125P5C8 protein whose sequence is encoded by (c) the cDNAs contained in the plasmids designated Escherichia coli DH5A 125P5C8PRO deposited with American Type Culture Collection as Accession No. PTA-3137;

(d) a polynucleotide that is fully complementary to a polynucleotide of any one of (a)-(c); and,

(e) a polynucleotide that selectively hybridizes under stringent conditions to a polynucleotide of (a)-(c).

Please cancel claim 15 and 20-22.

- 23. (Amended) A 125P5C8-related protein produced by a process comprising culturing a host cell that contains an expression vector comprising an 125P5C8 nucleotide, where T can be U, that comprises:
- (a) a polynucleotide having the sequence as shown in Figure 2 (SEQ ID NO: 1), from nucleotide residue number 1 through nucleotide residue number 2103; or,
- (b) a polynucleotide having the sequence as shown in Figure 2 (SEQ ID NO: 1), from nucleotide residue number 1 through nucleotide residue number 2100; or,
- (c) a polynucleotide having the sequence as shown in Figure 2 (SEQ ID NO: 1), from nucleotide residue number 1 through nucleotide residue number 2097; or
- (d) a polynucleotide of at least 10 bases of Figure 2 (SEQ ID NO: 1) that comprises the base at position 339;
- (e) a polynucleotide of at least 10 bases of Figure 2 (SEQ ID NO: 1) that comprises the base at position 1119;
- (f) a polynucleotide of at least \ 0 bases of Figure 2 (SEQ ID NO: 1) that comprises the base at position 2065;
- (g) a polynucleotide that selectively hybridizes under stringent conditions to a polynucleotide of (a)-(f);

wherein a range is understood to specifically disclose all whole unit positions thereof.